

Sortal Number: 09/998,284

CRF Errors Corrected by the STIC Systems Branch

2430 090
b103 OIPE

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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/998,284

DATE: 12/11/2001
TIME: 17:00:33

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\12112001\I998284.raw

#2.

3 <110> APPLICANT: POULSEN, et al.
 5 <120> TITLE OF INVENTION: COMPOSITION
 7 <130> FILE REFERENCE: 674509-2035
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/998,284
 C--> 9 <141> CURRENT FILING DATE: 2001-11-30
 9 <150> PRIOR APPLICATION NUMBER: PCT/IB00/00829
 10 <151> PRIOR FILING DATE: 2000-06-02
 12 <150> PRIOR APPLICATION NUMBER: GB 9913050.2
 13 <151> PRIOR FILING DATE: 1999-06-04
 15 <160> NUMBER OF SEQ ID NOS: 2
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 1644
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Chondrus crispus
 24 <400> SEQUENCE: 1
 25 atggctactt tgccacaaaa ggacccaggt tacattgtta ttgacgtcaa cgctggta 60
 27 ccagacaago ctgaccgaag attgccatcc atgaagcaag gtttcaacag aagatggatt 120
 29 ggtaccaaca tcgatttcgt ttacgtcggt tacactccac aagggtgcttg tactgctttg 180
 31 gacagagcta tggaaaagtg ttctccaggt accgtcagaa tcgtttctgg tggtaactgt 240
 33 tacgaagact tcgtttcga cgaatgtgtc aaggctatta tcaacgttac tggtttgggt 300
 35 gaatctggtt acgacgacga tagagggtac ttctgtcttt ccggtgacac caactgggg 360
 37 tccttcaaga ccttgttcag agaccacgggt agagtttgc caggtggttc ctgttactcc 420
 39 gtgggtttgg gtggtcacat tgcgggtgg ggtgacggta ttttggccag attgcacgg 480
 41 ttgccagtcg attggtttac cgggtgttggaa gttgtcggttta agccagtctt gaccgaagac 540
 43 tctgttctta agtacgttca caaggattcc gaaggttaacg acgggtgagtt gtttgggt 600
 45 cacactgggt gaggtggagg taacttcggt attatcacca aatactactt caagatttg 660
 47 ccaatgtctc caagagggtt catcgcttct aacttacact tctcttggga cggtttca 720
 49 agagatgcct tgcagattt gtgacttaag tacttcaagt tggtagatg tgattggaa 780
 51 aatactgttg gtaagttcca aatcttccac caagcagctg aagagttgt tatgtacttg 840
 53 tatacatctt actctaacga cggcgagaga gaagttgccc aagacagacata ctatcatttg 900
 55 gaggctgaca ttgaacagat ctacaaaaca tgcgagccta ccaaagctct tgggtgtcat 960
 57 gctgggttggg ctcccttccc ttttagaccc agaaagagac acacatccaa gacttcttat 1020
 59 atgcatgacg agactatggc ctacccttcc tacgcttga ctgagactat caacggttcc 1080
 61 ggtccataatc agagaggtaa gtacaagtct gcttacatga tcaaggactt tccagacttc 1140
 63 cagattgtatg ttatctggaa ataccttact gaggttccgt acgggttgc tagtggcgaa 1200
 65 atgaaggatg ctcttcttca gttgtatgt ttccgggttgg agattcacaa gtttgggttgg 1260
 67 gatgtactg cagttgtca gagagagtac atcatcaaac tgcagttacca gacatactgg 1320
 69 caggaagaag acaaggatgc agttaacttg aagtggatta gagacttttca cgaggagatg 1380
 71 tatgagcctt atgggtgttgc tccagaccct aacactcagg ttgagagtttcaaaaggtgtt 1440
 73 tttgagggt gctacttcaa ctaccctgat gttgacttga acaactggaa gaacggtaag 1500
 75 tatgggtgcct tggaaaccttta cttttgggt aacctgaaca gattgatcaa ggccaaatgg 1560
 77 ttgtgggatc ctaacgagat cttcacaaaac aaacagtcta tccctactaa acctcttaag 1620
 79 gagcctaagc agactaaata gtag 1644
 82 <210> SEQ ID NO: 2
 83 <211> LENGTH: 546
 84 <212> TYPE: PRT

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85 <213> ORGANISM: Chondrus crispus
87 <400> SEQUENCE: 2
89 Met Ala Thr Leu Pro Gln Lys Asp Pro Gly Tyr Ile Val Ile Asp Val
90 1 5 10 15
93 Asn Ala Gly Thr Pro Asp Lys Pro Asp Pro Arg Leu Pro Ser Met Lys
94 20 25 30
97 Gln Gly Phe Asn Arg Arg Trp Ile Gly Thr Asn Ile Asp Phe Val Tyr
98 35 40 45
101 Val Val Tyr Thr Pro Gln Gly Ala Cys Thr Ala Leu Asp Arg Ala Met
102 50 55 60
105 Glu Lys Cys Ser Pro Gly Thr Val Arg Ile Val Ser Gly Gly His Cys
106 65 70 75 80
109 Tyr Glu Asp Phe Val Phe Asp Glu Cys Val Lys Ala Ile Ile Asn Val
110 85 90 95
113 Thr Gly Leu Val Glu Ser Gly Tyr Asp Asp Asp Arg Gly Tyr Phe Val
114 100 105 110
117 Ser Ser Gly Asp Thr Asn Trp Gly Ser Phe Lys Thr Leu Phe Arg Asp
118 115 120 125
121 His Gly Arg Val Leu Pro Gly Gly Ser Cys Tyr Ser Val Gly Leu Gly
122 130 135 140
125 Gly His Ile Val Gly Gly Asp Gly Ile Leu Ala Arg Leu His Gly
126 145 150 155 160
129 Leu Pro Val Asp Trp Leu Ser Gly Val Glu Val Val Val Lys Pro Val
130 165 170 175
133 Leu Thr Glu Asp Ser Val Leu Lys Tyr Val His Lys Asp Ser Glu Gly
134 180 185 190
137 Asn Asp Gly Glu Leu Phe Trp Ala His Thr Gly Gly Gly Gly Asn
138 195 200 205
141 Phe Gly Ile Ile Thr Lys Tyr Tyr Phe Lys Asp Leu Pro Met Ser Pro
142 210 215 220
145 Arg Gly Val Ile Ala Ser Asn Leu His Phe Ser Trp Asp Gly Phe Thr
146 225 230 235 240
149 Arg Asp Ala Leu Gln Asp Leu Leu Thr Lys Tyr Phe Lys Leu Ala Arg
150 245 250 255
153 Cys Asp Trp Lys Asn Thr Val Gly Lys Phe Gln Ile Phe His Gln Ala
154 260 265 270
157 Ala Glu Glu Phe Val Met Tyr Leu Tyr Thr Ser Tyr Ser Asn Asp Ala
158 275 280 285
161 Glu Arg Glu Val Ala Gln Asp Arg His Tyr His Leu Glu Ala Asp Ile
162 290 295 300
165 Glu Gln Ile Tyr Lys Thr Cys Glu Pro Thr Lys Ala Leu Gly Gly His
166 305 310 315 320
169 Ala Gly Trp Ala Pro Phe Pro Val Arg Pro Arg Lys Arg His Thr Ser
170 325 330 335
173 Lys Thr Ser Tyr Met His Asp Glu Thr Met Asp Tyr Pro Phe Tyr Ala
174 340 345 350
177 Leu Thr Glu Thr Ile Asn Gly Ser Gly Pro Asn Gln Arg Gly Lys Tyr
178 355 360 365
181 Lys Ser Ala Tyr Met Ile Lys Asp Phe Pro Asp Phe Gln Ile Asp Val

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182 370 375 380
185 Ile Trp Lys Tyr Leu Thr Glu Val Pro Asp Gly Leu Thr Ser Ala Glu
186 385 390 395 400
189 Met Lys Asp Ala Leu Leu Gln Val Asp Met Phe Gly Gly Glu Ile His
190 405 410 415
193 Lys Val Val Trp Asp Ala Thr Ala Val Ala Gln Arg Glu Tyr Ile Ile
194 420 425 430
197 Lys Leu Gln Tyr Gln Thr Tyr Trp Gln Glu Glu Asp Lys Asp Ala Val
198 435 440 445
201 Asn Leu Lys Trp Ile Arg Asp Phe Tyr Glu Glu Met Tyr Glu Pro Tyr
202 450 455 460
205 Gly Gly Val Pro Asp Pro Asn Thr Gln Val Glu Ser Gly Lys Gly Val
206 465 470 475 480
209 Phe Glu Gly Cys Tyr Phe Asn Tyr Pro Asp Val Asp Leu Asn Asn Trp
210 485 490 495
213 Lys Asn Gly Lys Tyr Gly Ala Leu Glu Leu Tyr Phe Leu Gly Asn Leu
214 500 505 510
217 Asn Arg Leu Ile Lys Ala Lys Trp Leu Trp Asp Pro Asn Glu Ile Phe
218 515 520 525
221 Thr Asn Lys Gln Ser Ile Pro Thr Lys Pro Leu Lys Glu Pro Lys Gln
222 530 535 540
225 Thr Lys
226 545

VERIFICATION SUMMARY
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L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date